



		10	20	30	40	50	60
SEQ ID NO:2	HUMAN	MGIVEPGCGDMLTGTEPMPGSDEGRAPGADPQHRYFYYPEPGAQDADERRGGGSLGSPYPG					
SEQ ID NO:4	MOUSE	MGIVEPGCGDMLTGTEPMP-SDEGRGPGADQQHRFFFYYPEPGAQDPTDRRAGSSSLGTPYSG					
		10	20	30	40	50	
	CONS	MGIVEPGCGDMLTGTEPMP	SDEGR	PGAD	QHR	FYYPEPGAQD	RR G SLG PY G
		70	80	90	100	110	120
SEQ ID NO:2	HUMAN	GALVPAPPSRFLGAYAYPPRQAAGFPGAGESFPPPAEAGYQPGEGYAAPDPRAGLYPG					
SEQ ID NO:4	MOUSE	GALVPAAPGRFLGSAFYPPRAQVAGFPGPGEFFPPPAEAGYPPVDGYAPDPRAGLYPG					
		60	70	80	90	100	110
	CONS	GALVPA	P	RFLG	AYPPR	Q	AGFPG GE FPPPA AEGY P GY APDPRAGLYPG
		130	140	150	160	170	180
SEQ ID NO:2	HUMAN	PREDYALPAGLEVSGKLRVALNNHLLWSKFNQHQTEMIITKQGRRMFPFLSFTVAGLEPT					
SEQ ID NO:4	MOUSE	PREDYALPAGLEVSGKLRVALSNHLLWSKFNQHQTEMIITKQGRRMFPFLSFTVAGLEPT					
		120	130	140	150	160	170
	CONS	PREDYALPAGLEVSGKLRVAL	NHLLWSKFNQHQTEMIITKQGRRMFPFLSFTVAGLEPT				
		190	200	210	220	230	240
SEQ ID NO:2	HUMAN	SHYRMFVDVVLVDQHHWRYQSGKWVQCGKAEGSMPGNRLYVHPDSPNTGAHWMRQEVSFG					
SEQ ID NO:4	MOUSE	SHYRMFVDVVLVDQHHWRYQSGKWVQCGKAEGSMPGNRLYVHPDSPNTGAHWMRQEVSFG					
		180	190	200	210	220	230
	CONS	SHYRMFVDVVLVDQHHWRYQSGKWVQCGKAEGSMPGNRLYVHPDSPNTGAHWMRQEVSFG					
		250	260	270	280	290	300
SEQ ID NO:2	HUMAN	KLKLTNNKGASNNVTQMIVLQSLHKYQPRLHIVEVNDGEPEAACNASNTHIFFTQETQFI					
SEQ ID NO:4	MOUSE	KLKLTNNKGASNNVTQMIVLQSLHKYQPRLHIVEVNDGEPEAACASNTHVFTTQETQFI					
		240	250	260	270	280	290
	CONS	KLKLTNNKGASNNVTQMIVLQSLHKYQPRLHIVEVNDGEPEAAC	ASNTH	FTTQETQFI			

Fig. 1A

```

          310      320      330      340      350      360
SEQ ID NO:2 HUMAN AVTAYQNAEITQLKIDNNPFAKGFRENFESMYTSVDTSIPSPPGPNCQFLGGDHYSPLLP
          ::::::::::::::::::::::::::::::::::::::::::::::::::::::::::::::::::::
SEQ ID NO:4 MOUSE AVTAYQNAEITQLKIDNNPFAKGFRENFESMYASVDTSVPSPPGPNCQLLGDPFSPLLS
          300      310      320      330      340      350

```

CONS AVTAYQNAEITQLKIDNNPFAKGFREN FESMY SVDTS PSPPGPNCQ LGGD FSPLL

```

          370      380      390      400      410      420
SEQ ID NO:2 HUMAN NQYPVPSRFYPDLPGQAKDVVPQAYWLGAPRDHSYEAEFRAVSMKPAFLPSAPGPTMSYY
          :::::::::::::::::::::: : ::::::::::::::::::::::::::::::::::::::::::::
SEQ ID NO:4 MOUSE NQYPVPSRFYPDLPGQPKDMISQPYWLGTPREHSYEAEFRAVSMKPTLLPSAPGPTVPYY
          360      370      380      390      400      410

```

CONS NQYPVPSRFYPDLPGQ KD Q YWLG PR HSYEAEFRAVSMKP LPSAPGPT YY

```

          430      440      450      460      470      480
SEQ ID NO:2 HUMAN RGQEV LAPGAGWPVAPQYPPKMGPASWFRPMTLPMEPGGSEGRGPEDQGPPLVWTEI
          :::::::::::::::::::::::::::::::::::::::::::::::::::::: : : : : : :
SEQ ID NO:4 MOUSE RGQDV LAPGAGWPVAPQYPPKMSPAGWFRPMTLPM D PGLGSSEEQG----SSPSLWPEV
          420      430      440      450      460      470

```

CONS RGQ VLAPGAGWPVAPQYPPKM PA WFRPMTLPM PG G SE G P W E

```

          490      500      510      520      530
SEQ ID NO:2 HUMAN APIRPESSD SGLGEGDSKRRRVSPYPSSGDSSSPAGAPSPFDKEAEGQFYNTFPN
          . . . : : : : : : : : : : : : : : : : : : : : : : :
SEQ ID NO:4 MOUSE TSLQPEPSD SGLGEGDTKRRRISYPYPSSGDSSSPAGAPSPFDKETEGQFYNYFPN
          480      490      ↑↑↑ 500      510      520      530

```

CONS PE SDSGLGEGD KRRR SPYPSSGDSSSPAGAPSPFDKE EGQFYNYFPN

Fig. 1B

[illegible]

Fig. 1C

SEQ ID NO:1	HUMAN	CTCAACAACCACCTGTTGTGGTCCAAGTTTAATCAGCACCAGACAGAGATGATCATCACC
		430 440 450 460 470 480
SEQ ID NO:3	MOUSE	CTCAGCAACCACCTGTTGTGGTCCAAGTTCAACCAGCACCAGACAGAGATGATCATCACT
		420 430 440 450 460 470
SEQ ID NO:1	HUMAN	AAGCAGGGACGGCGGATGTTCCCATTCCTGTCATTTACTGTGGCCGGGCTGGAGCCCACC
		490 500 510 520 530 540
SEQ ID NO:3	MOUSE	AAGCAAGGACGGCGAATGTTCCCATTCCTGTCCTTACCCTGGCCGGGCTGGAGCCCACA
		480 490 500 510 520 530
SEQ ID NO:1	HUMAN	AGCCACTACAGGATGTTTGTGGACGTGGTCTTGGTGGACCAGCACCCTGGCGGTACCAG
		550 560 570 580 590 600
SEQ ID NO:3	MOUSE	AGCCATTACAGGATGTTTGTGGATGTGGTCTTGGTGGACCAGCACCCTGGCGGTACCAG
		540 550 560 570 580 590
SEQ ID NO:1	HUMAN	AGCGGCAAGTGGGTGCAGTGTGGAAAGGCCGAGGGCAGCATGCCAGGAAACCGCCTGTAC
		610 620 630 640 650 660
SEQ ID NO:3	MOUSE	AGCGGCAAGTGGGTGCAGTGTGGAAAGGCAGAAAGGCAGCATGCCAGGGAACCGCTTATAT
		600 610 620 630 640 650
SEQ ID NO:1	HUMAN	GTCCACCCGGACTCCCCAACACAGGAGCGCACTGGATGCGCCAGGAAGTTTCATTTGGG
		670 680 690 700 710 720
SEQ ID NO:3	MOUSE	GTCCACCCAGACTCCCCAACACCGGAGCCCACTGGATGCGCCAGGAAGTTTCATTTGGG
		660 670 680 690 700 710
SEQ ID NO:1	HUMAN	AAACTAAAGCTCACAAACAACAAGGGGGCGTCCAACAATGTGACCCAGATGATTGTGCTC
		730 740 750 760 770 780
SEQ ID NO:3	MOUSE	AAGCTAAAGCTCACCAACAACAAGGGGGCTTCCAACAATGTGACCCAGATGATCGTCCTG
		720 730 740 750 760 770
SEQ ID NO:1	HUMAN	CAGTCCCTCCATAAGTACCAGCCCCGGCTGCATATCGTTGAGGTGAACGACGGAGAGCCA
		790 800 810 820 830 840
SEQ ID NO:3	MOUSE	CAGTCTCTCCACAAGTACCAGCCCCGGCTGCACATCGTGGAGGTGAATGATGGAGAGCCA
		780 790 800 810 820 830

Fig. 1D

Fig. 1E

Fig. 1F

		10	20	30	40	50	60
<u>SEQ ID NO:2</u>	HUMAN	MGIVEPGCGDMLTGTEPMPGSDEGRAPGADPQHRYFYYPEPGAQDADERRGGGSLGSPYPG					
		:	:	:	:	:	:
<u>SEQ ID NO:4</u>	MOUSE	MGIVEPGCGDMLTGTEPMP-SDEGRGPGADQQHRFFYPEPGAQDPTDRRAGSSLGTPYSG					
		10	20	30	40	50	
	CONS	MGIVEPGCGDMLTGTEPMP	SDEGR	PGAD	QHR.FYYPEPGAQD	RR G	SLG PY G

		70	80	90	100	110	120
<u>SEQ ID NO:2</u>	HUMAN	GALVPAPPSRFLGAYAYPPRPQAAGFPGAGESFPPPADEGYQPGEGYAAPDPRAGLYPG					
		60	70	80	90	100	110
<u>SEQ ID NO:4</u>	MOUSE	GALVPAAPGRFLGSAFYPPRAQVAGFGPGGEFFPPPADEGYPPVDGYAPDPRAGLYPG					
	CONS	GALVPA P RFLG AYPPR Q AGFPG GE FPPPA AEGY P GY APDPRAGLYPG					

		130	140	150	160	170	180
<u>SEQ ID NO:2</u>	HUMAN	PREDYALPAGLEVSGKLRVALNNHLLWSKFNQHQTEMIITKQGRRMFPFLSFTVAGLEPT ::					
<u>SEQ ID NO:4</u>	MOUSE	PREDYALPAGLEVSGKLRVALSNHLLWSKFNQHQTEMIITKQGRRMFPFLSFTVAGLEPT 120 130 140 150 160 170					
	CONS	PREDYALPAGLEVSGKLRVAL NHLLWSKFNQHQTEMIITKQGRRMFPFLSFTVAGLEPT					

		190	200	210	220	230	240
<u>SEQ ID NO:2</u>	HUMAN	SHYRMFVDVVLVDQHHWRYQSGKWVQCGKAEGSMPGNRLYVHPDSPNTGAHWMRQEV SFG					
		::					
<u>SEQ ID NO:4</u>	MOUSE	SHYRMFVDVVLVDQHHWRYQSGKWVQCGKAEGSMPGNRLYVHPDSPNTGAHWMRQEV SFG					
		180	190	200	210	220	230
	CONS	SHYRMFVDVVLVDQHHWRYQSGKWVQCGKAEGSMPGNRLYVHPDSPNTGAHWMRQEV SFG					

		250	260	270	280	290	300
<u>SEQ ID NO:2</u>	HUMAN	KLKLTNNKGASNNVTQMIVLQSLHKYQPR	LHIVEVNDGEPEAACNASNTH	FTTFQETQFI			
		::::::::::::::::::::::::::::::::	::::::::::::::::::::::::::::::::	::::::::::::::::::::::::::::::::	::::::::::::::::::::::::::::::::	::::::::::::::::::::::::::::::::	::::::::::::::::::::::::::::::::
<u>SEQ ID NO:4</u>	MOUSE	KLKLTNNKGASNNVTQMIVLQSLHKYQPR	LHIVEVNDGEPEAAC	SASNTHVFTTFQETQFI			
		240	250	260	270	280	290
	CONS	KLKLTNNKGASNNVTQMIVLQSLHKYQPR	LHIVEVNDGEPEAAC	ASNTH	FTFOETOFI		

Fig. 1A

SEQ ID NO:2

SEQ ID NO:4

CONS AVTAYQNAEITQLKIDNNPFAKGFRENFESMY SVDTS PSPPGPNCQ LGGD FSPLL

SEQ ID NO:2

SEQ ID NO:4

CONS NQYPVPSRFYPLPGQ KD Q YWLG PR HSYEAEFRAVSMKP LPSAPGPT YY

SEQ ID NO:2

SEQ ID NO: 4

CONS RGQ VLAPGAGWPVAPQYPPKM PA WFRPMRTLPM PG G SE G P W E

SEQ ID NO:2

SEQ ID NO:4

CONS PE SDSGLGED KRRR SPYPSSGDSSSPAGAPSPFDKE EGQFYNYFPN

Fig. 1B ~~Fig. 1A (continued)~~

SEQ ID NO:1	HUMAN	10	20	30	40	50	60
		ATGGGCATCGTGGAGCCGGGTTGCGGAGACATGCTGACGGGCACCGAGCCGATGCCGGGG					
SEQ ID NO:3	MOUSE	ATGGGCATCGTGGAGCCGGGCTGCGGAGACATGCTGACCGGCACCGAGCCGATGCC---G					
		10	20	30	40	50	
SEQ ID NO:1	HUMAN	70	80	90	100	110	120
		AGCGACGAGGGCCGGGCGCCTGGCGCCGACCCGACGACCGCTACTTCTACCCGGAGCCG					
SEQ ID NO:3	MOUSE	AGTGACGAGGGCCGGGGGCCCGGAGCGGACCAACAGCATCGTTTCTTCTATCCCGAGCCG					
		60	70	80	90	100	110
SEQ ID NO:1	HUMAN	130	140	150	160	170	180
		GGCGCGCAGGACGCGGACGAGCGTCGCGGGGGCGGCAGCCTGGGGTCTCCCTACCCGGGG					
SEQ ID NO:3	MOUSE	GGCGCACAGGACCCGACCGATCGCCGCGCAGGTAGCAGCCTGGGGACGCCCTACTCTGGG					
		120	130	140	150	160	170
SEQ ID NO:1	HUMAN	190	200	210	220	230	240
		GGCGCCTTGGTGCCCGCCCCGCGAGCCGCTTCCTTGGAGCCTACGCCTACCCGCCGCGA					
SEQ ID NO:3	MOUSE	GCGCCCTGGTGCCCTGCCGCGCCGGGTCGCTTCCTTGGATCCTTCGCCTACCCGCCCGG					
		180	190	200	210	220	230
SEQ ID NO:1	HUMAN	250	260	270	280	290	300
		CCCCAGGCGGCCGGCTTCCCCGGCGCGGGCGAGTCCTTCCCGCCGCCCGCGGACGCCGAG					
SEQ ID NO:3	MOUSE	GCTCAGGTGGCTGGCTTTCGGGCGCTGGCGAGTTCTTCCCGCCGCCCGGGGTGCGGAG					
		240	250	260	270	280	290
SEQ ID NO:1	HUMAN	310	320	330	340	350	360
		GGCTACCAGCCGGGCGAGGGCTACGCCGCCCCGGACCCGCGCGCCGGGCTCTACCCGGGG					
SEQ ID NO:3	MOUSE	GGCTACCCGCCCGTGGATGGCTACCCTGCCCTGACCCGCGCGCGGGGCTCTACCCAGGG					
		300	310	320	330	340	350
SEQ ID NO:1	HUMAN	370	380	390	400	410	420
		CCGCGTGAGGACTACGCGCTACCCGCGGGACTGGAGGTGTCGGGGAAACTGAGGGTCGCG					
SEQ ID NO:3	MOUSE	CCGCGCGAGGACTACGCATTGCCCGCGGGGTGGAGGTGTCGGGAAGCTGAGAGTCGCG					
		360	370	380	390	400	410

Fig. 1C Fig. 1B

SEQ ID NO:1	HUMAN	CTCAACAACCACTGTTGTGGTCCAAGTTTAATCAGCACCAGACAGAGATGATCATCACC
		430 440 450 460 470 480
SEQ ID NO:3	MOUSE	CTCAGCAACCACTGTTGTGGTCCAAGTTCAACCAGCACCAGACAGAGATGATCATCACT
		420 430 440 450 460 470
SEQ ID NO:1	HUMAN	AAGCAGGGACGGCGGATGTTCCCATTCCTGTCAATTTACTGTGGCCGGGCTGGAGCCCACC
		490 500 510 520 530 540
SEQ ID NO:3	MOUSE	AAGCAAGGACGGCGAATGTTCCCATTCCTGTCTTCACCGTGGCCGGGCTGGAGCCCACA
		480 490 500 510 520 530
SEQ ID NO:1	HUMAN	AGCCACTACAGGATGTTTGTGGACGTGGTCTTGGTGGACCAGCACCAGTGGCGGTACCAG
		550 560 570 580 590 600
SEQ ID NO:3	MOUSE	AGCCATTACAGGATGTTTGTGGATGTGGTCTTGGTGGACCAGCACCAGTGGCGGTACCAG
		540 550 560 570 580 590
SEQ ID NO:1	HUMAN	AGCGGCAAGTGGGTGCAGTGTGGAAAGGCCGAGGGCAGCATGCCAGGAAACCGCCTGTAC
		610 620 630 640 650 660
SEQ ID NO:3	MOUSE	AGCGGCAAGTGGGTGCAGTGTGGAAAGGCAGAAGGCAGCATGCCAGGGAACCGCTTATAT
		600 610 620 630 640 650
SEQ ID NO:1	HUMAN	GTCCACCCGGA CTCCCCAACACAGGAGCGCACTGGATGCGCCAGGAAGTTTCATTGGG
		670 680 690 700 710 720
SEQ ID NO:3	MOUSE	GTCCACCCAGACTCCCCAACACCGGAGCCCACTGGATGCGCCAGGAAGTTTCATTGGG
		660 670 680 690 700 710
SEQ ID NO:1	HUMAN	AAACTAAAGCTCACAAACAACAAGGGGGCGTCCAACAATGTGACCCAGATGATTGTGCTC
		730 740 750 760 770 780
SEQ ID NO:3	MOUSE	AAGCTAAAGCTCACCAACAACAAGGGGGCTTCCAACAATGTGACCCAGATGATCGTCCTG
		720 730 740 750 760 770
SEQ ID NO:1	HUMAN	CAGTCCCTCCATAAGTACCAGCCCCGGCTGCATATCGTTGAGGTGAACGACGGAGAGCCA
		790 800 810 820 830 840
SEQ ID NO:3	MOUSE	CAGTCTCTCCACAAGTACCAGCCCCGGCTGCACATCGTGGAGGTGAATGATGGAGAGCCA
		780 790 800 810 820 830

Fig. 1D ~~Fig. 1B~~ (continued)

		850	860	870	880	890	900
<u>SEQ ID NO:1</u>	HUMAN	GAGGCAGCCTGCAACGCTTCCAACACGCATATCTTTACTTTCCAAGAAACCCAGTTCATT					
		::::	::::	::::	::::	::::	::::
<u>SEQ ID NO:3</u>	MOUSE	GAGGCTGCCTGCAGTGCTTCTAACACACACGTCTTTACTTTCCAAGAGACCCAGTTCATT					
		840	850	860	870	880	890

		910	920	930	940	950	960
<u>SEQ ID NO:1</u>	HUMAN	GCCGTGACTGCCTACCAGAATGCCGAGATTACTCAGCTGAAAATTGATAATAACCCCTTT					
		::	::::	::::	::::	::::	::::
<u>SEQ ID NO:3</u>	MOUSE	GCAGTGACTGCCTACCAGAACGCAGAGATCACTCAGCTGAAAATCGACAACAACCCCTTT					
		900	910	920	930	940	950

		970	980	990	1000	1010	1020
<u>SEQ ID NO:1</u>	HUMAN	GCCAAAGGATTCCGGGAGAACCTTTGAGTCCATGTACACATCTGTTGACACCAGCATCCCC					
		::::	::::	::::	::::	::::	::::
<u>SEQ ID NO:3</u>	MOUSE	GCCAAAGGATTCCGGGAGAACCTTTGAGTCCATGTACGCATCTGTTGATACGAGTGTCCCC					
		960	970	980	990	1000	1010

		1030	1040	1050	1060	1070	1080
<u>SEQ ID NO:1</u>	HUMAN	TCCCCGCCTGGACCCAACCTGTCAATTCCTTGGGGGAGATCACTACTCTCCTCTCCTACCC					
		::	::	::::	::::	::::	::::
<u>SEQ ID NO:3</u>	MOUSE	TCGCCACCTGGACCCAACCTGTCAACTGCTTGGGGGAGACCCCTTCTCACCTCTTCTATCC					
		1020	1030	1040	1050	1060	1070

		1090	1100	1110	1120	1130	1140
<u>SEQ ID NO:1</u>	HUMAN	AACCAGTATCCTGTTCCCAGCCGCTTCTACCCCGACCTTCCTGGCCAGGCCAAGGATGTG					
		::::	::::	::::	::::	::::	::::
<u>SEQ ID NO:3</u>	MOUSE	AACCAGTATCCTGTTCCCAGCCGTTTCTACCCCGACCTTCAGGCCAGCCCAAGGATATG					
		1080	1090	1100	1110	1120	1130

		1150	1160	1170	1180	1190	1200
<u>SEQ ID NO:1</u>	HUMAN	GTTCCCCAGGCTTACTGGCTGGGGGCCCCCGGGACCACAGCTATGAGGCTGAGTTTCGA					
		:	:	::::	::::	::::	::::
<u>SEQ ID NO:3</u>	MOUSE	ATCTCACAGCCTTACTGGCTGGGGACACCTCGGGAACACAGTTATGAAGCGGAGTTCCGA					
		1140	1150	1160	1170	1180	1190

		1210	1220	1230	1240	1250	1260
<u>SEQ ID NO:1</u>	HUMAN	GCAGTCAGCATGAAGCCTGCATTCTTGCCCTCTGCCCCTGGGCCACCATGTCCTACTAC					
		::	::	::::	::::	::::	::::
<u>SEQ ID NO:3</u>	MOUSE	GCTGTGAGCATGAAGCCCACTCTACCTCTGCCCCGGGGCCCACTGTGCCCTACTAC					
		1200	1210	1220	1230	1240	1250

Fig. 1E ~~Fig. 1B~~ (continued)

SEQ ID NO:1 HUMAN 1270 1280 1290 1300 1310 1320
CGAGGCCAGGAGGTCTGGCACCTGGAGCTGGCTGGCCTGTGGCACCCCAGTACCCCTCCC
:: ::::: :: ::::: ::::: ::::: ::::: ::::: ::::: ::::: ::::: ::::: :::::
SEQ ID NO:3 MOUSE 1260 1270 1280 1290 1300 1310
CGGGGCCAAGACGTCTGGCGCCTGGAGCTGGTTGGCCCGTGGCCCTCAATACCCGCC

SEQ ID NO:1 HUMAN 1330 1340 1350 1360 1370 1380
AAGATGGGCCCCGGCCAGCTGGTTCCGCCCTATGCGGACTCTGCCCATGGAACCCGGCCCT
::: ::::: ::::: ::::: ::::: ::::: ::::: ::::: ::::: ::::: :::::
SEQ ID NO:3 MOUSE 1320 1330 1340 1350 1360 1370
AAGATGAGCCCAGCTGGCTGGTTCCGGCCCATGCGAACTCTGCCCATGGACCCGGGCCCTG

SEQ ID NO:1 HUMAN 1390 1400 1410 1420 1430 1440
GGAGGCTCAGAGGGACGGGACCAGAGGACCAGGGTCCCCCTTGGTGTGGACTGAGATT
::: ::::: ::::: ::::: ::::: ::::: ::::: ::::: ::::: ::::: :::::
SEQ ID NO:3 MOUSE 1380 1390 1400 1410 1420
GGATCCTCAGAGGAACAGGGCTCCT-----CCCCCTCGCTGTGGCCTGAGGTC

SEQ ID NO:1 HUMAN 1450 1460 1470 1480 1490 1500
GCCCCATCCGGCCGGAATCCAGTGATTTCAGGACTGGGCGAAGGAGACTCTAAGAGGAGG
::: ::::: ::::: ::::: ::::: ::::: ::::: ::::: ::::: ::::: :::::
SEQ ID NO:3 MOUSE 1430 1440 1450 1460 1470 1480
ACCTCCCTCCAGCCGAGCCAGCGACTCAGGACTAGGCGAAGGAGACACTAAGAGGAGG

SEQ ID NO:1 HUMAN 1510 1520 1530 1540 1550 1560
CGCGTGTCCCCCTATCCTTCCAGTGGTGACAGCTCCTCCCCTGCTGGGGCCCCCTTCTCCT
: : ::::: ::::: ::::: ::::: ::::: ::::: ::::: ::::: ::::: :::::
SEQ ID NO:3 MOUSE 1490 1500 1510 1520 1530 1540
AGGATATCCCCCTATCCTTCCAGTGGCGACAGCTCCTCTCCCGCTGGGGCCCCCTTCTCCT

SEQ ID NO:1 HUMAN 1570 1580 1590 1600
TTTGATAAGGAAGCTGAAGGACAGTTTATAACTATTTTCCCAACTGA
::: ::::: ::::: ::::: ::::: ::::: ::::: ::::: ::::: :::::
SEQ ID NO:3 MOUSE 1550 1560 1570 1580 1590
TTTGATAAGGAAACCGAAGGCCAGTTTATAATTATTTTCCCAACTGA

Fig. 1F ~~Fig. 1B~~ (continued)